	Changed the margins in cases where the sequence text was "wrapped" down to the next line.
	Edited a format error in the Current Application Data section, specifically:
	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other
	Added the mandatory heading and subheadings for "Current Application Data".
	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
	Changed the spelling of a mandatory field (the headings or subheadings), specifically:
	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
l	nserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
	Inserted colons after headings/subheadings. Headings edited included:
	Deleted extra, invalid, headings used by an applicant, specifically:
	Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of page numbers throughout text; other invalid text, such as
	Inserted mandatory headings, specifically:
	Corrected an obvious error in the response, specifically:
	Edited identifiers where upper case is used but lower case is required, or vice versa.
	Corrected an error in the Number of Sequences field, specifically:
	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
	Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (errodue to a Patentin bug). Sequences corrected:
	Other:

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.



PCT10

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1A1F: 11 18 1
                      RAW SEQUENCE LISTING
                                                            0.1MF: 0.8:45:4
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      5 <110> APPRICANT: Skubite, Keith M.
              Skubitz, Amy F.N.
      % K120~ TitlE (F HWENTION: Poptides Capable of Modulatin: The Fination of CD66 (CEACAM
Family
              Members
     ll kisch file beffbende: 28210-011
     1 - <140 - CURPENT APPLICATION NUMBER: 10/069,605
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     2\times \times 1510^\circ PRIOR FILING DATE: 2000-08-25 21\times 160^\circ NUMBER OF SEQ ID NOS: 200
     20 -170. SOFTWARE: Fatentin version 3.1
     3 + 4210: 3EQ 1E NO: 1
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+ 1. FEATURE:
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65 Wal lie lys der Aspile: Wal Ashi Hip Hip Ala Thr Hip Hin
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89 KAISU SEQ II NO: 5
90 kölir LENGTH: 14
91 KICH TYPF: EFT
92 K2035 CEGANION: Abtoin that Sequence
94 02010 FEATURE:
93 (2039 (THER INFORMATION: Cynthetic Pertide
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99 Pro Glu The Min Asy The The Tyr Leu Trp Trp Ile Ash Ash.
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108 K22G> FEATURE:
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136 MASS OTUTE INVESTMENTION: Eventheric Fortide
138 <400> SEQUENCE: 8
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144 KU167 SEQ 11 N : 9
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140 FIRE ORGANISM: Artificial Sequence
149 HILLS FEATURE:
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THE REPORT OF THE MEDICAL CON-
194 Ala Ala Ser Alb Bro Bro Ala Sub Tyr Ser Try Double Ast
18 + 4.1 + 18, 11 N : 1
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16. .1. < IYFE: FFT</pre>
161 - 2152 "REANIOM: Artificial Cequence
163 KOVIN FRATURE:
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166 -416> CEQUENTE:
188 for the Ash By The The Win Bin Ser Thr Bin Wil Ten Fhe
  12 k2105 JEG ID NO: 11
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IMG KILZA IYER: BAT
175 Kliby OFBANIOM: Artifficial Sequence
179 R2105 FEATURE:
Translage Commission 1000: Synthetic Pertine
180 K400 % SEQUENTS: II
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ZZU P220> FEATURE:
001 (223) OTHER INFORMATION: Cynthetic Leptide
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. H. H. IHH. B. WILLIMI, Art. formal, Deguence
184 KILL K BRAIMBE:
. He - . . - - THEE THE EMATE WE Dynthetic Teptide
137 -416 - 089,78078: 13
1.59 Old Ard Mit live Led Der Bln. Bly Ash Thr Thr led Der Ile
..40 1
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144 KILLS IFN TH: 14
246 KLIZ - THIE: FET
.46 <218 OFGANISM: Arti: Poial Sequence
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282 I
285 <210: SEQ ID NO: 19
286 <2112 LENGTH: 14
UST «2020 TYPE: FRT
288 (2003) OCHADIME ARTIEL (al Chigue en
ZAU KYSU. ALAMANIA
291 <223: OTHER INFORMATION: Synthetic Peptide
193 <4001 SEQUENCE: 19
195 Leu val His Ash Leu Ero Gin Asp Ero Arg Gly Tyr Ash Trp
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tob Kulla LENGIH: 14
BOI RØIDH TYFE: FRI
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RAW SEQUENCE LISTING DATE: II . . . 10MB: 19:45:4 EA ENT APELL ATT N: US/10/069,605 : A:\PTO.AMC.txt :: : N:\CRF4\11132002\J069605.raw + 4 *... * FFAITHE: Following the COR RMATION: Synthetic Express * Proposition of the Application Application Application Applications · 1 : :14 - 111 - 1EM31H: 14 318 KUINA TYFE: ERI 316 VIII - CEGANUM: Artificial Sequence 318 - 22 - FRATURE: RIB WOOK OFFER INFORMATION: Synthetic Feptide MAI W4 - PRANCENCE: UI 323 Val ile lys ieu Ash Leu Met Ser Sid Gld Val ihr Sig Sin 3.27 3/9 KOIN KIYEE: EED HBO KLOP KRAANINM: Artificial Sequence dda FVIT - FEAITER: 333 klubs OTHER INFORMATION: Synthetic Feptide 335 K4CLS SEQUENCE: 22 337 Fig. 31d Thr. 4th Ash Thr Thr Tyr Led Trp Trp Val Ash 3ty 338 1 341 ×011 × 2EQ 11 % : 13 342 KD11 - LENGTH: 14 343 - KLOP TYFF: 1FT 344 <213 < OFGANIOM: Artificial Sequence 346 KELOF FEATURE: 347 (213) OTHER INFOFMATION: Synthetic Peptide 349 <400> SEQUENCE: 13 001 Fre Cer Asy Thr Tyr Tyr Nis Ala Cly Val Aon Leu Aon Leu 352 1 355 <210> SEQ ID NO: 24 356 <011> LENGTH: 14 357 <212> TYPE: PHT 358 KAISH ORGANISM: Artificial Sequence 3€0 <220> FEATURE: 361 <223> OTHER INFORMATION: Synthetic Peptide 363 ×4000 SEQUENTE: 24 M65 Ala Ala Ser Arm Tro Tro Cor Clm Tyr Ser Trp Cor Tal Acm 166 369 <210> SEQ ID NO: 25 376 KO11> LENGTH: 14 *71 <010> TYPF: FPT 302 k.13 (CRGANISM: Artificial Sequence :T4 < NOTE: FEATURE:

377 + 446 × SEQUEDICE: 15

*TE - LI: - CTHER INFORMATION: Symthetic legitide

37% Ser Val Ash Sly thr the Sin Gir Tyr Thr Sin Lys Leu Fre

 VERIFICATION SUMMARY
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